

[illegible][illegible]



[illegible]

with the following motifs? (A1AM 157) (A1AM 157).

$$\begin{aligned} \mathbf{A} \cdot \mathbf{A}^T \mathbf{M} &= \mathbf{I}^T \\ \mathbf{M} \mathbf{M}^T &= \mathbf{I} \end{aligned} \quad \text{if } \mathbf{M} = \begin{pmatrix} \cos \theta & \sin \theta \\ -\sin \theta & \cos \theta \end{pmatrix}.$$

Euprimates: Eutheria; Placentalia; Primates; Catarrhini; Hominoidea; Homo.

10 908 PROBABLY H2 901

11 RESULT 1

12 AT2.FOVIN

13 ID AT2.FOVIN STANDARD PRG: 1205.AA

14 15 OCT 2001 (Oct. 40, 2001)

15 16 OCT 2001 (Oct. 40, 2001) Last sequence update

16 17 JUN 2002 (Oct. 41, 2002) Last annotation update

17 18 OCT 2002 (Oct. 41, 2002) Last annotation update

18 19 OCT 2002 (Oct. 41, 2002) Last annotation update

19 20 OCT 2002 (Oct. 41, 2002) Last annotation update

20 21 OCT 2002 (Oct. 41, 2002) Last annotation update

21 22 OCT 2002 (Oct. 41, 2002) Last annotation update

22 23 OCT 2002 (Oct. 41, 2002) Last annotation update

23 24 OCT 2002 (Oct. 41, 2002) Last annotation update

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75 76 OCT 2002 (Oct. 41, 2002) Last annotation update

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79 DBL: X06389, CAA5255.1

80 MEMO: M2.301

81 INTERP: 100007622: Disintegrin

82 INTERP: 100007622: pep_M2H_p1000

83 INTERP: 100007622: Repolysin

84 INTERP: 100007622: TSP

85 INTERP: 100007622: TSP

86 INTERP: 100007622: TSP

87 INTERP: 100007622: TSP

88 INTERP: 100007622: TSP

89 INTERP: 100007622: TSP

90 INTERP: 100007622: TSP

91 INTERP: 100007622: TSP

92 INTERP: 100007622: TSP

93 INTERP: 100007622: TSP

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Search time 26.2002, 16:25:40 : Search time 26.2002

(without a query file)
4682015 Million cell updates/sec

Post-processed: 1 MAWA/ALGAL/MIMIC/.../FOFOSAYEPHMOCKPOCH 1104

Post-processed: 1 MAWA/ALGAL/MIMIC/.../FOFOSAYEPHMOCKPOCH 1104

Total number of hits satisfying chosen parameters: 283224

Maximum hit size: 200000000

Post-processed: Maximum Match 0%
Maximum Match 100%
Listed first 45 summaries

Post-processed: 1 MAWA/ALGAL/MIMIC/.../FOFOSAYEPHMOCKPOCH 1104

Post-processed: 1 MAWA/ALGAL/MIMIC/.../FOFOSAYEPHMOCKROCH 1104

SUMMARIES

Rank	Score	Match	Length	ID	Description
1	144.5	28.5	2	T18517	procollagen N-endo
2	144.5	22.1	2	T121371	hypothetical prote
3	129.5	21.2	2	T00017	gene ADAMS-1 prot
4	129.5	18.0	2	T00355	hypothetical prote
5	85.5	14.0	2	T47158	hypothetical prote
6	84.5	14.5	2	C89114	protein C17C3.6a
7	82.4	14.5	2	T134395	hypothetical prote
8	79.5	11.6	2	T18856	hypothetical prote
9	74.4	10.9	2	T00260	hypothetical prote
10	74.4	9.6	2	T16892	hypothetical prote
11	74.4	8.7	2	T22545	hypothetical prote
12	74.4	7.7	2	T15976	hypothetical prote
13	74.4	7.7	2	T15976	hypothetical prote
14	74.4	7.7	2	T15976	hypothetical prote
15	74.4	7.7	2	T15976	hypothetical prote
16	74.4	7.7	2	T15976	hypothetical prote
17	74.4	7.7	2	T15976	hypothetical prote
18	74.4	7.7	2	T15976	hypothetical prote
19	74.4	7.7	2	T15976	hypothetical prote
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21	74.4	7.7	2	T15976	hypothetical prote
22	74.4	7.7	2	T15976	hypothetical prote
23	74.4	7.7	2	T15976	hypothetical prote
24	74.4	7.7	2	T15976	hypothetical prote
25	74.4	7.7	2	T15976	hypothetical prote
26	74.4	7.7	2	T15976	hypothetical prote
27	74.4	7.7	2	T15976	hypothetical prote
28	74.4	7.7	2	T15976	hypothetical prote
29	74.4	7.7	2	T15976	hypothetical prote

40	109	1.1	2	HYSA
41	109	1.1	2	HYSA
42	109	1.1	2	HYSA
43	109	1.1	2	HYSA
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100	109	1.1	2	HYSA

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844 030620 N. A. V. K. A.

827 GOWRIE, J. J. N. and S. E. ARKIN, 1971, p. 191

[illegible]

RESULT 4

Hypothetical $\mu_1 = 0.5$ and $\mu_2 = 0.5$
 C-Species: 100 and 100

CONCLUSION: THE

1. *Escherichia coli* 167 (167) 167

A. K. KOTCHERIKOVA, *Department of Chemistry, Moscow State University, Moscow, U.S.S.R.*

A-Accession: 1000
A-Species: *Pseudomonas*
A-Material: Protein; DNA

A; Residues: 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100, 101, 102, 103, 104, 105, 106, 107, 108, 109, 110, 111, 112, 113, 114, 115, 116, 117, 118, 119, 120, 121, 122, 123, 124, 125, 126, 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138, 139, 140, 141, 142, 143, 144, 145, 146, 147, 148, 149, 150, 151, 152, 153, 154, 155, 156, 157, 158, 159, 160, 161, 162, 163, 164, 165, 166, 167, 168, 169, 170, 171, 172, 173, 174, 175, 176, 177, 178, 179, 180, 181, 182, 183, 184, 185, 186, 187, 188, 189, 190, 191, 192, 193, 194, 195, 196, 197, 198, 199, 200, 201, 202, 203, 204, 205, 206, 207, 208, 209, 210, 211, 212, 213, 214, 215, 216, 217, 218, 219, 220, 221, 222, 223, 224, 225, 226, 227, 228, 229, 230, 231, 232, 233, 234, 235, 236, 237, 238, 239, 240, 241, 242, 243, 244, 245, 246, 247, 248, 249, 250, 251, 252, 253, 254, 255, 256, 257, 258, 259, 260, 261, 262, 263, 264, 265, 266, 267, 268, 269, 270, 271, 272, 273, 274, 275, 276, 277, 278, 279, 280, 281, 282, 283, 284, 285, 286, 287, 288, 289, 290, 291, 292, 293, 294, 295, 296, 297, 298, 299, 300, 301, 302, 303, 304, 305, 306, 307, 308, 309, 310, 311, 312, 313, 314, 315, 316, 317, 318, 319, 320, 321, 322, 323, 324, 325, 326, 327, 328, 329, 330, 331, 332, 333, 334, 335, 336, 337, 338, 339, 340, 341, 342, 343, 344, 345, 346, 347, 348, 349, 350, 351, 352, 353, 354, 355, 356, 357, 358, 359, 360, 361, 362, 363, 364, 365, 366, 367, 368, 369, 370, 371, 372, 373, 374, 375, 376, 377, 378, 379, 380, 381, 382, 383, 384, 385, 386, 387, 388, 389, 390, 391, 392, 393, 394, 395, 396, 397, 398, 399, 400, 401, 402, 403, 404, 405, 406, 407, 408, 409, 410, 411, 412, 413, 414, 415, 416, 417, 418, 419, 420, 421, 422, 423, 424, 425, 426, 427, 428, 429, 430, 431, 432, 433, 434, 435, 436, 437, 438, 439, 440, 441, 442, 443, 444, 445, 446, 447, 448, 449, 450, 451, 452, 453, 454, 455, 456, 457, 458, 459, 460, 461, 462, 463, 464, 465, 466, 467, 468, 469, 470, 471, 472, 473, 474, 475, 476, 477, 478, 479, 480, 481, 482, 483, 484, 485, 486, 487, 488, 489, 490, 491, 492, 493, 494, 495, 496, 497, 498, 499, 500, 501, 502, 503, 504, 505, 506, 507, 508, 509, 510, 511, 512, 513, 514, 515, 516, 517, 518, 519, 520, 521, 522, 523, 524, 525, 526, 527, 528, 529, 530, 531, 532, 533, 534, 535, 536, 537, 538, 539, 540, 541, 542, 543, 544, 545, 546, 547, 548, 549, 550, 551, 552, 553, 554, 555, 556, 557, 558, 559, 560, 561, 562, 563, 564, 565, 566, 567, 568, 569, 570, 571, 572, 573, 574, 575, 576, 577, 578, 579, 580, 581, 582, 583, 584, 585, 586, 587, 588, 589, 590, 591, 592, 593, 594, 595, 596, 597, 598, 599, 600, 601, 602, 603, 604, 605, 606, 607, 608, 609, 610, 611, 612, 613, 614, 615, 616, 617, 618, 619, 620, 621, 622, 623, 624, 625, 626, 627, 628, 629, 630, 631, 632, 633, 634, 635, 636, 637, 638, 639, 640, 641, 642, 643, 644, 645, 646, 647, 648, 649, 650, 651, 652, 653, 654, 655, 656, 657, 658, 659, 660, 661, 662, 663, 664, 665, 666, 667, 668, 669, 670, 671, 672, 673, 674, 675, 676, 677, 678, 679, 680, 681, 682, 683, 684, 685, 686, 687, 688, 689, 690, 691, 692, 693, 694, 695, 696, 697, 698, 699, 700, 701, 702, 703, 704, 705, 706, 707, 708, 709, 710, 711, 712, 713, 714, 715, 716, 717, 718, 719, 720, 721, 722, 723, 724, 725, 726, 727, 728, 729, 730, 731, 732, 733, 734, 735, 736, 737, 738, 739, 740, 741, 742, 743, 744, 745, 746, 747, 748, 749, 750, 751, 752, 753, 754, 755, 756, 757, 758, 759, 760, 761, 762, 763, 764, 765, 766, 767, 768, 769, 770, 771, 772, 773, 774, 775, 776, 777, 778, 779, 780, 781, 782, 783, 784, 785, 786, 787, 788, 789, 790, 791, 792, 793, 794, 795, 796, 797, 798, 799, 800, 801, 802, 803, 804, 805, 806, 807, 808, 809, 810, 811, 812, 813, 814, 815, 816, 817, 818, 819, 820, 821, 822, 823, 824, 825, 826, 827, 828, 829, 830, 831, 832, 833, 834, 835, 836, 837, 838, 839, 8

A: Experimental $\alpha = 0.05$
C: Bonferroni's

Agilene: K1A40188

F:519 575/Enthail. 11 sp. coll. 1960-1961

COPY	MATERIALS
BEST COPY AVAILABLE	FBI - NEW YORK

[illegible]

40 EXHIBIT EN - ATTORNEY GENERAL

[illegible][illegible]

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1

[illegible][illegible]

Quality Match:	9.68	Score	585.5	DB 2:	Length	860	
Host Local Similarity:	24.48	Pred. No.	1.7e-31				
Matches:	449	Mismatches:	127	Totals:	307	Gaps:	53

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Search completed : 09/04/2015 12:00:10
Job time : 34 sec.



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XX At isolated nucleic acid molecule encoding a novel human
 F1 metallopeptidase family member useful for diagnosing neurological
 F2 disorders and inflammatory response
 XX

ES Claim 9: Fig. 1A-E: 15ppp: English.

XX The invention relates to an isolated nucleic acid molecule designated
 3442a, which encodes a novel metallopeptidase family member. The nucleic
 acid encodes a metallopeptidase which may be useful for treating or
 preventing a neurological disorder (pain related disorder), or conduct
 or may also be of use for determining presence or absence of genetic
 alteration for disease diagnosis. My state is useful as novel diagnostic
 markers and therapeutic agents for bone disorders, immune disorders,
 cardiovascular disorders, liver disorders, viral diseases, skeletal
 muscle disorders, reproductive disorders, skin disorders, kidney
 disorders and adipose and thyroid disorders. The agent that modulates the
 activity or expression of the nucleic acid is also used in the invention.
 XX The present sequence represents an additional human 3442a protein sequence.

XX Sequence: 1444 AA

XX Match: 94.4%, Score: 57421, ID: 24, Length: 1044

XX Match: 1044, Score: 11, Mismatch: 0, Indels: 0, Gaps: 0

XX 1201 HETEROLOGOUSLY EXPRESSED IN A BACTERIAL HOST AND PURIFIED AS A
 1202 1203 1204 1205 1206 1207 1208 1209 1210 1211 1212 1213 1214 1215 1216 1217 1218 1219 1220 1221 1222 1223 1224 1225 1226 1227 1228 1229 1230 1231 1232 1233 1234 1235 1236 1237 1238 1239 1240 1241 1242 1243 1244 1245 1246 1247 1248 1249 1250 1251 1252 1253 1254 1255 1256 1257 1258 1259 1260 1261 1262 1263 1264 1265 1266 1267 1268 1269 1270 1271 1272 1273 1274 1275 1276 1277 1278 1279 1280 1281 1282 1283 1284 1285 1286 1287 1288 1289 1290 1291 1292 1293 1294 1295 1296 1297 1298 1299 1300 1301 1302 1303 1304 1305 1306 1307 1308 1309 1310 1311 1312 1313 1314 1315 1316 1317 1318 1319 1320 1321 1322 1323 1324 1325 1326 1327 1328 1329 1330 1331 1332 1333 1334 1335 1336 1337 1338 1339 1340 1341 1342 1343 1344 1345 1346 1347 1348 1349 1350 1351 1352 1353 1354 1355 1356 1357 1358 1359 1360 1361 1362 1363 1364 1365 1366 1367 1368 1369 1370 1371 1372 1373 1374 1375 1376 1377 1378 1379 1380 1381 1382 1383 1384 1385 1386 1387 1388 1389 1390 1391 1392 1393 1394 1395 1396 1397 1398 1399 1400 1401 1402 1403 1404 1405 1406 1407 1408 1409 1410 1411 1412 1413 1414 1415 1416 1417 1418 1419 1420 1421 1422 1423 1424 1425 1426 1427 1428 1429 1430 1431 1432 1433 1434 1435 1436 1437 1438 1439 1440 1441 1442 1443 1444 1445 1446 1447 1448 1449 1450 1451 1452 1453 1454 1455 1456 1457 1458 1459 1460 1461 1462 1463 1464 1465 1466 1467 1468 1469 1470 1471 1472 1473 1474 1475 1476 1477 1478 1479 1480 1481 1482 1483 1484 1485 1486 1487 1488 1489 1490 1491 1492 1493 1494 1495 1496 1497 1498 1499 1500 1501 1502 1503 1504 1505 1506 1507 1508 1509 1510 1511 1512 1513 1514 1515 1516 1517 1518 1519 1520 1521 1522 1523 1524 1525 1526 1527 1528 1529 1530 1531 1532 1533 1534 1535 1536 1537 1538 1539 1540 1541 1542 1543 1544 1545 1546 1547 1548 1549 1550 1551 1552 1553 1554 1555 1556 1557 1558 1559 1560 1561 1562 1563 1564 1565 1566 1567 1568 1569 1570 1571 1572 1573 1574 1575 1576 1577 1578 1579 1580 1581 1582 1583 1584 1585 1586 1587 1588 1589 1590 1591 1592 1593 1594 1595 1596 1597 1598 1599 1600 1601 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Xgapext	0.5
Ygapext	0.5
Fgapext	7.0
Delcext	7.0

14530575366 Residuals

ing chosen parameters:

Age Group	Total (%)	18-29 (%)	30-49 (%)	50-69 (%)	70+ (%)
18-29	88	92	85	82	78
30-49	85	82	88	85	80
50-69	82	78	85	88	82
70+	78	75	82	85	88

100

Part 45 Summaries

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at 2002 Oct 16 17:00:00 MAT
-STRAT=10 -N=1000000
OUT=6 -OUTL=1-7
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of results from the

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SUMMARY

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09-04746 (1-1114) * 2000 12 20 42

24b. VALIN, A. and L. G. M. 1979. The effect of the temperature of the water on the growth of the European sea bass, *Morone labrax* L. *Journal of the European Association of Fish Pathologists*, 19: 111-114.

[illegible]

coronary heart failure; myocardial infarction; cerebral ischemia;
 neurodegenerative disease; high infection's disease; peripheral neuropathy;
 cerebral amyloid angiopathy; amyotrophic lateral sclerosis; multiple
 multiple sclerosis; cerebral angiogenesis; normal injury; depression;
 mental degeneration; abnormal wound healing; burns; diabetic shock; ds.

Key: (C) Copyright 1999
 139, 414
 /Product: "AluMTS-E"

EP149900-A1
 11-01-2001

26-APR-2001 ZOLTEP 030818.

26-APR-2001 ZOLTEP 030818.

(PFI) Pfizer Inc.

Buckley, L. Mitchell, P. Walsh, N. Wachtman, T.

WPI, 2002 01/01/03.

PFI, 2002 01/01/03.

NOTE: This document contains information about a distribution and
 metalloproteinase containing thrombospondin domain, and AluMTS-E2
 polypeptide used for treating cerebral ischemia, osteoporosis,
 cerebral, allergies and asthma.

claim 1: Page 16 17 18pp (end of)

This sequence encodes AluMTS-E which is a disintegrin and metallo-
 protease containing a thrombospondin domain. AluMTS-E2 is a splice
 variant of AluMTS-E. The AluMTS-E DNA is useful for diagnosing a
 disease or susceptibility to a disease in a subject relative to
 expression of activity of AluMTS-E in a subject. AluMTS-E and the
 splice variant may be used in the treatment of osteoarthritis and
 rheumatoid arthritis, inflammatory bowel disease, Crohn's disease,
 emphysema, acute respiratory distress syndrome, asthma, chronic
 obstructive pulmonary disease, Alzheimer's disease, organ transplant
 rejection and rejection, leukemia, allergy, cancer, tissue ulcerations,
 psychosis, peritoneal disease, epidermal cysts, bullous, osteoporosis,
 osteoporosis of artificial joint implants, atherosclerosis, aortic
 aneurysm, congestive heart failure, myocardial infarction, stroke,
 cerebral ischemia, head trauma, spinal cord injury, neurodegenerative
 disease, autoimmune disorders, Huntington's disease, Parkinson's
 disease, multiple sclerosis, periph. neuropathy, pain, cerebral
 amyloid angiopathy, amyotrophic lateral sclerosis, amyotrophic
 lateral sclerosis, multiple sclerosis, ocular angiogenesis, cerebral
 injury, mental degeneration, abnormal wound healing, burns, infection
 or diabetic shock.

Sequence 1726-60, 710-60, 1204-60, 1292-60, 1311-60, 1311-60

Alignment Scores:

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AATKZ1572

22. 11. 2017 (11.11.2017)

Public Affairs Officer, Ministry of Foreign Affairs, Tokyo

kinetics of the reaction of the metal protease, M01S1, with a substrate, and metal protease type M01S1.

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THE UNIVERSITY OF CHICAGO

Local Communities

322

7. *Effect of "a distilling" and met-enkephalinase (ADM)*

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carriers, arthritis and rheumatoid arthritis

The present section is devoted to the construction of a separable Hilbert space $\mathcal{H}(\mathcal{W})$ associated to \mathcal{W} .

1. The first part of the document is a list of references. The references are listed in a standard format, with the author's name, the title of the work, and the publisher. The references are as follows:

1. J. H. Van Veen, *The History of the Netherlands*, 1910, 1911, 1912, 1913, 1914, 1915, 1916, 1917, 1918, 1919, 1920, 1921, 1922, 1923, 1924, 1925, 1926, 1927, 1928, 1929, 1930, 1931, 1932, 1933, 1934, 1935, 1936, 1937, 1938, 1939, 1940, 1941, 1942, 1943, 1944, 1945, 1946, 1947, 1948, 1949, 1950, 1951, 1952, 1953, 1954, 1955, 1956, 1957, 1958, 1959, 1960, 1961, 1962, 1963, 1964, 1965, 1966, 1967, 1968, 1969, 1970, 1971, 1972, 1973, 1974, 1975, 1976, 1977, 1978, 1979, 1980, 1981, 1982, 1983, 1984, 1985, 1986, 1987, 1988, 1989, 1990, 1991, 1992, 1993, 1994, 1995, 1996, 1997, 1998, 1999, 2000, 2001, 2002, 2003, 2004, 2005, 2006, 2007, 2008, 2009, 2010, 2011, 2012, 2013, 2014, 2015, 2016, 2017, 2018, 2019, 2020, 2021, 2022, 2023, 2024, 2025, 2026, 2027, 2028, 2029, 2030, 2031, 2032, 2033, 2034, 2035, 2036, 2037, 2038, 2039, 2040, 2041, 2042, 2043, 2044, 2045, 2046, 2047, 2048, 2049, 2050, 2051, 2052, 2053, 2054, 2055, 2056, 2057, 2058, 2059, 2060, 2061, 2062, 2063, 2064, 2065, 2066, 2067, 2068, 2069, 2070, 2071, 2072, 2073, 2074, 2075, 2076, 2077, 2078, 2079, 2080, 2081, 2082, 2083, 2084, 2085, 2086, 2087, 2088, 2089, 2090, 2091, 2092, 2093, 2094, 2095, 2096, 2097, 2098, 2099, 2100, 2101, 2102, 2103, 2104, 2105, 2106, 2107, 2108, 2109, 2110, 2111, 2112, 2113, 2114, 2115, 2116, 2117, 2118, 2119, 2120, 2121, 2122, 2123, 2124, 2125, 2126, 2127, 2128, 2129, 2130, 2131, 2132, 2133, 2134, 2135, 2136, 2137, 2138, 2139, 2140, 2141, 2142, 2143, 2144, 2145, 2146, 2147, 2148, 2149, 2150, 2151, 2152, 2153, 2154, 2155, 2156, 2157, 2158, 2159, 2160, 2161, 2162, 2163, 2164, 2165, 2166, 2167, 2168, 2169, 2170, 2171, 2172, 2173, 2174, 2175, 2176, 2177, 2178, 2179, 2180, 2181, 2182, 2183, 2184, 2185, 2186, 2187, 2188, 2189, 2190, 2191, 2192, 2193, 2194, 2195, 2196, 2197, 2198, 2199, 2200, 2201, 2202, 2203, 2204, 2205, 2206, 2207, 2208, 2209, 2210, 2211, 2212, 2213, 2214, 2215, 2216, 2217, 2218, 2219, 2220, 2221, 2222, 2223, 2224, 2225, 2226, 2227, 2228, 2229, 2230, 2231, 2232, 2233, 2234, 2235, 2236, 2237, 2238, 2239, 2240, 2241, 2242, 2243, 2244, 2245, 2246, 2247, 2248, 2249, 2250, 2251, 2252, 2253, 2254, 2255, 2256, 2257, 2258, 2259, 2260, 2261, 2262, 2263, 2264, 2265, 2266, 2267, 2268, 2269, 2270, 2271, 2272, 2273, 2274, 2275, 2276, 2277, 2278, 2279, 2280, 2281, 2282, 2283, 2284, 2285, 2286, 2287, 2288, 2289, 2290, 2291, 2292, 2293, 2294, 2295, 2296, 2297, 2298, 2299, 2300, 2301, 2302, 2303, 2304, 2305, 2306, 2307, 2308, 2309, 2310, 2311, 2312, 2313, 2314, 2315, 2316, 2317, 2318, 2319, 2320, 2321, 2322, 2323, 2324, 2325, 2326, 2327, 2328, 2329, 2330, 2331, 2332, 2333, 2334, 2335, 2336, 2337, 2338, 2339, 2340, 2341, 2342, 2343, 2344, 2345, 2346, 2347, 2348, 2349, 2350, 2351, 2352, 2353, 2354, 2355, 2356, 2357, 2358, 2359, 2360, 2361, 2362, 2363, 2364, 2365, 2366, 2367, 2368, 2369, 2370, 2371, 2372, 2373, 2374, 2375, 2376, 2377, 2378, 2379, 2380, 2381, 2382, 2383, 2384, 2385, 2386, 2387, 2388, 2389, 2390, 2391, 2392, 2393, 2394, 2395, 2396, 2397, 2398, 2399, 2400, 2401, 2402, 2403, 2404, 2405, 2406, 2407, 2408, 2409, 2410, 2411, 2412, 2413, 2414, 2415, 2416, 2417, 2418, 2419, 2420, 2421, 2422, 2423, 2424, 2425, 2426, 2427, 2428, 2429, 2430, 2431, 2432, 2433, 2434, 2435, 2436, 2437, 2438, 2439, 2440, 2441, 2442, 2443, 2444, 2445, 2446, 2447, 2448, 2449, 2450, 2451, 2452, 2453, 2454, 2455, 2456, 2457, 2458, 2459, 2460, 2461, 2462, 2463, 2464, 2465, 2466, 2467, 2468, 2469, 2470, 2471, 2472, 2473, 2474, 2475, 2476, 2477, 2478, 2479, 2480, 2481, 2482, 2483, 2484, 2485, 2486, 2487, 2488, 2489, 2490, 2491, 2492, 2493, 2494, 2495, 2496, 2497, 2498, 2499, 2500, 2501, 2502, 2503, 2504, 2505, 2506, 2507, 2508, 2509, 2510, 2511, 2512, 2513, 2514, 2515, 2516, 2517, 2518, 2519, 2520, 2521, 2522, 2523, 2524, 2525, 2526, 2527, 2528, 2529, 2530, 2531, 2532, 2533, 2534, 2535, 2536, 2537, 2538, 2539, 2540, 2541, 2542, 2543, 2544, 2545, 2546, 2547, 2548, 2549, 2550, 2551, 2552, 2553, 2554, 2555, 2556, 2557, 2558, 2559, 2560, 2561, 2562, 2563, 2564, 2565, 2566, 2567, 2568, 2569, 2570, 2571, 2572, 2573, 2574, 2575, 2576, 2577, 2578,

$$E_{\text{eff}}(t) = N(t) \cdot \sum_i E_i(t) \quad [eV]$$

1. The first part of the document is a list of names and their corresponding page numbers. The names are listed in a single column, and the page numbers are listed in a single column to the right of the names. The names are: "The first part of the document is a list of names and their corresponding page numbers."

Met (1): (y_1, z_1) Index: 1

1

Age Group	Percentage
18-24	10%
25-34	15%
35-44	20%
45-54	25%
55-64	20%
65-74	15%
75-84	10%
85-94	5%
95-104	5%

1000

1

[illegible][illegible]

[illegible][illegible]

RESULT 4
ASS-97.74

[illegible]

[illegible]

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487	488	489	490	491	492	493	494	495	496	497
498	499	500	501	502	503	504	505	506	507	508
509	510	511	512	513	514	515				

[illegible]

[illegible][illegible]

[illegible]

421 AIDAHI S. I. *Chem. Abstr.* 1954, 48, 11434a [AIDAHI S. I., *Yuki Kagaku*, 1953, 11, 149].



[illegible]

[illegible]

AN ROJON
WEEKLY IN
KIEW-TOO
SINGAPORE
STATION
TELEPHONE
HONG KONG

Mammalia: Monotremata: Ornithorhina; Vertebrata: Euteleostomi:
 Mammalia: Euphrasia: Primates: Catarrhini: Hominoidea: Homo.
 1 (1998: 1-644)
 REFERENCE
 AUTHOR'S ADDRESS:

Department of Psychology, University of Maryland,
 3501 Research Building, College Park, MD 20742
 E-mail: jacob@psych.umd.edu
 Received 10 June 1997
 Accepted 10 June 1998
 Copyright © 1998 John Wiley & Sons, Ltd.
J. Forecast. **17**, 103–114 (1998)

[illegible]

/-tattcsmc=Homo sapiens"
 /-Xref="axon:9638"
 /-IMAGE=254543"
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 /-loc_start="TetA"
 /-loc_end=DH10B (phase 2;start)"
 /-loc_organt="Thymus; Eukaryota; Vector: pTZ19-pac
 (pharmacia) with a modified polylinker Site.1; Not I;
 Site.2; Eco RI; 1st strand cDNA was primed with a Not I
 /-loc(h)= Primer 15'
 /-cDNA=ATGTGAGTCACCAAGGGGAAATTTTITTTTTTTT 3';
 A 16mer stranded cDNA was ligated to Eco RI adaptors
 (Pharmacia), digested with Not I and cleaved into the Not I
 and Eco RI sites of the modified pTZ19 vector. Library
 was titrated one round of normalization. library
 constructed by Genio Soares and M. Fatima Bonafide.

\mathcal{A}
 \mathcal{B}
 \mathcal{C}
 \mathcal{D}
 \mathcal{E}
 \mathcal{F}
 \mathcal{G}
 \mathcal{H}
 \mathcal{I}
 \mathcal{J}
 \mathcal{K}
 \mathcal{L}
 \mathcal{M}
 \mathcal{N}
 \mathcal{O}
 \mathcal{P}
 \mathcal{Q}
 \mathcal{R}
 \mathcal{S}
 \mathcal{T}
 \mathcal{U}
 \mathcal{V}
 \mathcal{W}
 \mathcal{X}
 \mathcal{Y}
 \mathcal{Z}

[illegible][illegible][illegible][illegible]

	97
Dd	97

07 886 PROSELYTIC VIOLENCE IN THE
08 457 COMMUNITARIAN

OF
1904
LIBRARY
OF THE
AMERICAN
MUSEUM OF
NATURAL HISTORY
1000
AMERICAN
MUSEUM OF
NATURAL HISTORY
1000

[illegible]

346 *PROCEEDINGS OF THE 1997 ANNUAL MEETING OF THE*
AMERICAN SOCIETY OF CLIMATE AND WATER RESOURCES
11-13 DECEMBER 1997

466 HISABAYASHI, YAMAGUCHI, AND KAWABATA

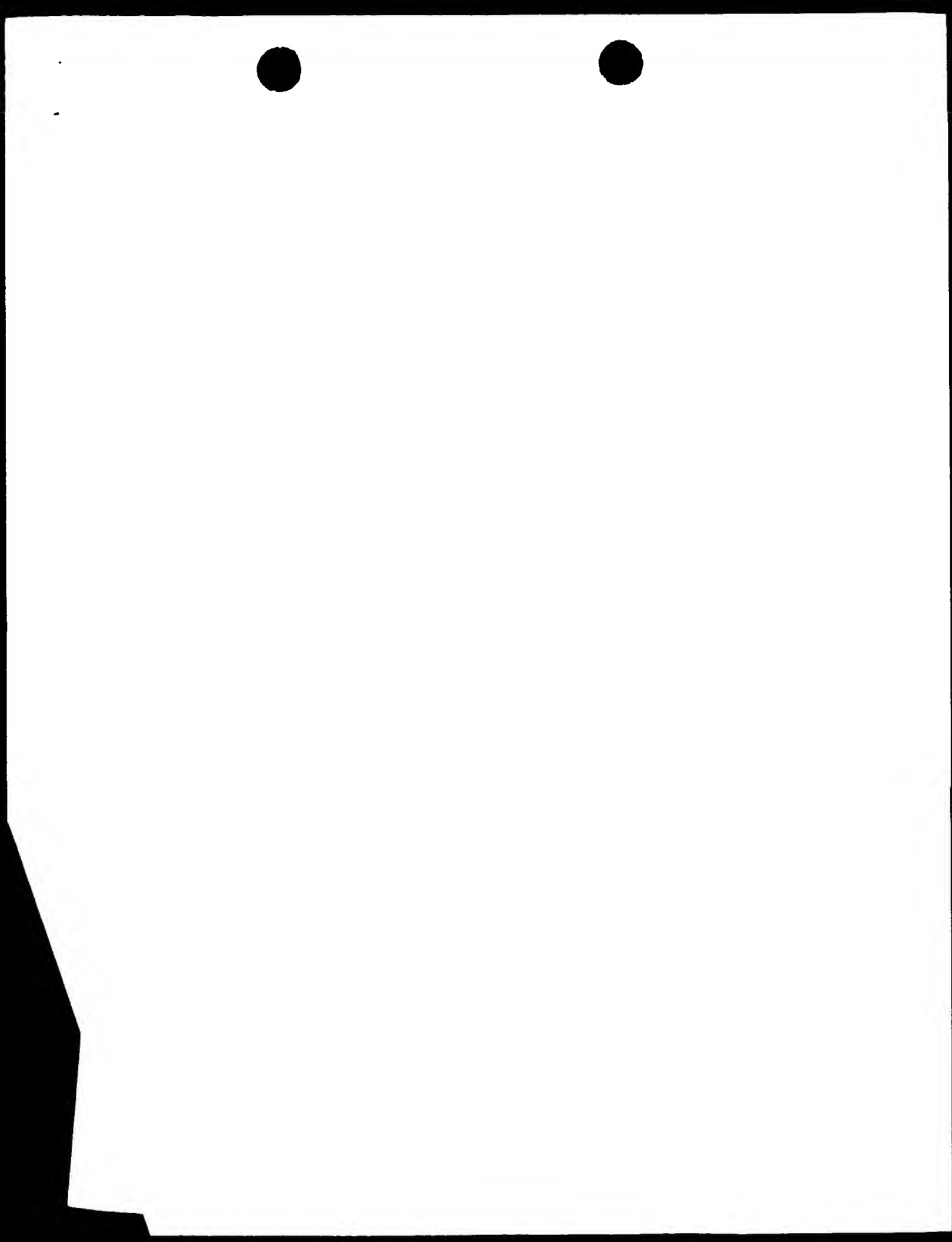
1. $\text{C}_2\text{H}_5\text{Br}$, $\text{C}_2\text{H}_5\text{I}$, $\text{C}_2\text{H}_5\text{Cl}$, $\text{C}_2\text{H}_5\text{F}$, $\text{C}_2\text{H}_5\text{Br}$, $\text{C}_2\text{H}_5\text{I}$, $\text{C}_2\text{H}_5\text{Cl}$, $\text{C}_2\text{H}_5\text{F}$	986	45
2. $\text{C}_2\text{H}_5\text{Br}$, $\text{C}_2\text{H}_5\text{I}$, $\text{C}_2\text{H}_5\text{Cl}$, $\text{C}_2\text{H}_5\text{F}$, $\text{C}_2\text{H}_5\text{Br}$, $\text{C}_2\text{H}_5\text{I}$, $\text{C}_2\text{H}_5\text{Cl}$, $\text{C}_2\text{H}_5\text{F}$	417	46

106 157 158 A
QY 1006 Arq11.....

1076 ANSON, A. AND CHAN, C. H. H. 1990. The

[illegible]

Statistical significance was assessed using the Fisher's exact test. A P value of ≤ 0.05 was considered significant.



TEACHES

COMMENT:

The distribution MDC clone distribution information can be found through the L.M.A.C.E. Consortium Web Site at www.lmac.org. This page has the following problems: frame shifted.

Lodgen/Vallieres

[illegible]

[illegible]

[illegible][illegible]



CM profile with a local frame_plus_F2b model

479f. 426 M) : 1103: (92)

100-104240-2

Xgapex	6.5
Ygapex	0.5
Egapex	7.0
Delex	7.0

197730502 res:1.2.5

1. The following chosen parameters: 71064...

Abstract:

M.I.T. in March 08
 M.I.T. in March 1009

List and list 45 summaries

1. $\frac{1}{2}$
 2. $\frac{1}{4}$
 3. $\frac{1}{8}$
 4. $\frac{1}{16}$
 5. $\frac{1}{32}$
 6. $\frac{1}{64}$
 7. $\frac{1}{128}$
 8. $\frac{1}{256}$
 9. $\frac{1}{512}$
 10. $\frac{1}{1024}$
 11. $\frac{1}{2048}$
 12. $\frac{1}{4096}$
 13. $\frac{1}{8192}$
 14. $\frac{1}{16384}$
 15. $\frac{1}{32768}$
 16. $\frac{1}{65536}$
 17. $\frac{1}{131072}$
 18. $\frac{1}{262144}$
 19. $\frac{1}{524288}$
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 21. $\frac{1}{2097152}$
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 40. $\frac{1}{1099511627776}$
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 43. $\frac{1}{8796093022208}$
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 11

[illegible]

total class : 0
: b. school Applications_NA : *

[illegible]

fixed. Now, as the number of results predicted by response to have a higher probability than or equal to the score of the results being predicted is observed by analysis of the total score distribution.

SUMMARY

Index	Match	Length	AppID	Description
1	5.42	16	08-09-858-001-10	Sequence 10, App1
2	5.42	10	08-09-858-001-10	Sequence 8, App1
3	5.42	10	08-09-858-001-11	Sequence 11, App1
4	5.41	10	08-09-858-001-3	Sequence 3, App1

1026)

[illegible]

ANNEX

[illegible]

[illegible][illegible]

[illegible][illegible]

Aluminum Structure	4.76e-14	Leaves:	3207
Fuel No.		Matches:	1038
Sieve	5.97e-09	Connections:	
Percent Surface	66.74%	Mistakes:	1
Best case simulation	96.65%	Inputs:	44
Geometry Material	94.05%	Gaps:	2
Density	19		

[illegible][illegible]

[illegible][illegible]

